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EP-A- 0 038 182**EP-A- 0 079 739****EP-A- 0 206 733****EP-A- 0 236 210****EP-A- 0 308 381**

PROC. NATL. ACAD. SCI. U.S.A. vol.79, Jan.
1982, p.71-75, Washington, US; A.Dugaiczky
et al: "Nucleotide sequence and the encoded
amino acids of human serum albumin
mRNA"

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EUR. J. BIOCHEM. vol.98, no.2 1979,p.477-485,
Berlin; R.T.A. Macgillivray et al.
:"Biosynthesis of Bovine plasma Proteins in
a cell free System"

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DescriptionFIELD OF THE INVENTION

5 This invention relates to a method for preparing foreign protein in yeast using an expression recombinant DNA comprising DNA encoding the serum albumin signal peptide adjacent to DNA encoding the foreign protein.

BACKGROUND OF THE INVENTION

10 In the production of specific proteins in a recombinant host by recombinant DNA technology, there are many advantages to having the host express and secrete the desired protein. That is, when a desired protein is expressed directly within the host cell, if there is any toxicity which inhibits growth or compromises the survival of the host cell, this toxicity can be avoided by the secretion of the protein. Even
 15 when there is no toxicity, as the protein accumulates in the host cell, it may inhibit the host cell growth. This, too, can be avoided by secretory expression. In addition, systems which accumulate protein in the host cell may also denature it, rendering it insoluble. This problem also can be avoided by secretory expression. Moreover, when commercially producing protein by recombinant DNA technology in a system which accumulates the desired protein intracellularly, it is necessary to destroy the cell in order to refine the
 20 protein, and it must be purified from the debris of the cellular destruction. This makes it difficult to obtain a protein of high purity. On the other hand, when producing a protein by a secretory expression system the protein only must be harvested from the culture broth, minimizing the problem of separating impurities derived from the recombinant host. This is a great advantage. Finally, most protein undergoes some modification, such as the addition of a sugar moiety, the formation of a disulfide bond, activation by limited
 25 hydrolysis of the inert proprotein, phosphorylation of specific amino acids, or carboxylation before activation. Some of these functions are performed by the themselves, and several of these modifications take place in the process of secretion. Therefore, a system which produces protein by secretory expression, as compared to a system which accumulates protein intracellularly, may be expected to generate proteins having a structure and function much close to the native protein.

30 Somethings are known about the properties of the signal peptide, and the characteristics of its amino acid sequence seem to be as follows. There are many basic amino acids near the N-terminal, and there are many polar amino acids near the portion which is digested by signal peptidase on the C-terminal side, while a sequence hydrophobic amino acids fill in the space between these two areas. The basic amino acids near the N-terminal interact with the phospholipids on the internal surface of the cell membrane, and the
 35 sequence of hydrophobic amino acids in the middle region plays an important role in passing the protein through the cell membrane. The polar amino acids at the C-terminal are believed to play some role in recognition during digestion by signal peptidase. These characteristics are extremely similar from pro-caryotes to higher animals, suggesting a common mechanism for protein secretion. (M.S. Briggs and L.M. Gierasch, Adv. Protein Chem., 38, 109-180 (1986); G. von Heijne, EMBO J., 3, 2315-2318 (1984)).

40 Human serum albumin is encoded on the gene as a prepro type protein (see Japanese Patent Application (OPI) No. 29985/87 (the term OPI used herein means an unexamined published application.) or EP-A-206733; A. Dugaiczky et al. Proc. Natl. Acad. Sci. USA, 79, 71-75 (1982)). The DNA and amino acid sequence in the vicinity of the N-terminal of mature human serum albumin beginning from the signal peptide essential for secretion are shown in Table 1 below.

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Table 1

	-----	Signal peptide	-----		Propeptide		
-18				-1	-6		-1 +1

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerAlaTyrSerArgGlyValPheArgArgAspAlaHis
ATGAAGTGGGTAAACCTTTATTTCCTTCCTTTCTCTTTAGCTGGCTTATTCACGGGGTGTGTTCGTCGAGATGCACAC

The signal peptide, composed of 18 amino acid is removed at the time of secretion. The propeptide, composed of 6 amino acids, is removed by processing, and mature human serum albumin, composed of 585 amino acids, and having an N-terminal amino acid sequence of Asp-Ala-His-Lys-Ser, is obtained.

Since yeast secrete less extracellular proteases and are capable of adding sugar moieties to its secret, yeast is excellent for the secretory expression of foreign proteins.

Several cases of signal peptides which contributes to the secretory expression in cells other than yeast, but which also function in yeast, have been reported. Examples include the secretory expression in yeast of human lysozyme using the chicken lysozyme signal peptide (Jigami, BIOINDUSTRY, 4, 117-123 (1987)), secretory expression in yeast of thaumatin using the signal peptide for plant protein thaumatin (L. Edens, I. Bom, A.M. Ledeboer, J. Maat, M.Y. Toonen, C. Visser and C.T. Verrips, Cell, 37, 629-633 (1984)), and secretory expression in yeast of human interferon using the signal peptide for human interferon- α (R.A. Hitzeman, D.W. Leung, L.J. Perry, W.J. Kohr, H.L. Levine and D.V. Doeddel, Science, 219, 620-625 (1983)).

The truth is, however, that the signal peptide contributing to secretory expression in cells other than yeast does not always function in yeast.

SUMMARY OF THE INVENTION

Therefore, a primary object of this invention is to provide a method for expressing and secreting foreign protein efficiently in yeast, the signal peptide gene functionable in yeast for secretory expression, the vector to be used in this method, and the transformant transformed by this vector.

The above-described object of the present invention has been met in one embodiment by a method for preparing foreign protein comprising expressing and secreting said foreign protein by yeast transformed by a recombinant DNA comprising the serum albumin signal peptide gene adjacent to the gene of said foreign protein. In a second embodiment, the present invention relates to a serum albumin signal peptide gene and derivatives thereof. In a third embodiment, the present invention relates to a recombinant DNA for transforming yeast comprising DNA encoding the serum albumin signal peptide adjacent to DNA encoding a foreign protein. In a forth embodiment, the present invention relates to a strain of yeast transformed by a recombinant DNA comprising DNA coding for the serum albumin signal peptide adjacent to DNA encoding a foreign protein.

BRIEF DESCRIPTION OF THE DRAWINGS

Fig. 1 shows the procedure for making pGAL12 from pGAL11 possessing the GAL1, 10 promoters.

Fig. 2 shows the procedure for making pPT1, containing only the pho5 terminator, from pAP5 and pUC9 containing the entire pho5 gene.

Fig. 3 shows the procedure for making pPT2 from pJDB207.

Fig. 4 and Fig. 5 show the restriction enzyme map of pGX401 containing the prepro human serum albumin gene.

Fig. 6 shows the procedure for making pHSA2, containing the human serum albumin gene C-terminal side from pGX401 and pUC19.

Fig. 7 shows the procedure for making pHSA1, containing the human serum albumin gene N-terminal side, from pGX401 and pUC19.

Fig. 8 shows the procedure for making pNH001, containing the signal peptide gene and the mature human serum albumin gene, from pHSA1, pHSA2 and the synthesized signal peptide gene.

Fig. 9 shows the procedure for making pNH007, containing the GAL1 promoter, signal peptide gene and mature human serum albumin gene, from pNH001 and pGAL12.

Fig. 10 shows the procedure for making pNH008, containing the GAL1 promoter, signal peptide gene, mature human serum albumin gene and pho5 terminator, from pNH007 and pPT2.

DETAILED DESCRIPTION OF THE INVENTION

The recombinant DNA of this invention comprises the serum albumin signal peptide gene, the foreign protein gene, a promoter, a terminator, and the plasmid DNA or chromosome DNA.

The origin of the serum albumin signal peptide gene is not specifically defined as long as it is derived from mammals. Practically, human-derived, rat-derived and bovine-derived preparations can be used.

Examples of the amino acid sequences of such signal peptides are known to include;
Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe Ser Ser Ala Tyr Ser derived from humans;
Met Lys Trp Val Thr Phe Leu Leu Leu Leu Phe Ile Ser Gly Ser Ala Phe Ser derived from rats; and
Met Lys Trp Val Thr Phe Ile Ser Leu Leu Leu Leu Phe Ser Ser Ala Tyr Ser derived from cows.

However, preferably, the human serum albumin signal peptide gene is used and the 2nd amino acid and the last five amino acids can be changed by Y and Xs as the following sequence.

Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu

Phe X₅ X₄ X₃ X₂ X₁

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wherein Y represents Lys, Arg or His and preferably represents Lys; X₅ represents Ala, Pro or Ser; X₄ represents Lys, Gly or Ser; X₃ represents Ala, Val or Cys and preferably represents Val or Cys; X₂ represents Tyr, Trp or Ser; and X₁ represents Ser, Ala or Gly and preferably represents Ala or Gly. Preferable examples of amino acid sequences of the signal peptides are shown in Table 2 below.

Table 2

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Sequence No.	Y	X ₅	X ₄	X ₃	X ₂	X ₁
Sequence 1	Lys	Ser	Ser	Val	Tyr	Ala
Sequence 2	Lys	Ala	Lys	Val	Ser	Ala
Sequence 3	Lys	Pro	Gly	Cys	Trp	Ala
Sequence 4	Lys	Pro	Gly	Val	Trp	Ala

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The serum albumin signal peptide gene may possess a DNA sequence which can be expressed by the amino acid sequence shown above, and one example is having the following DNA sequence.

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ATGAAGTGGGTAACCTTTATTTCCCTT

CTTTTCTCTTTAGCTCGGCTTATTC

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Preferable codons corresponding to each amino acid are set forth below.

Ala: GCT or GCC, Cys: TGT, Asp: GAC,

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Glu: GAA, Phe: TTC, Gly: GGT,

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His: GAC, Ile: ATT or ATC, Lys: AAG,

Leu: TTG, Met: ATG, Asn: AAC,

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Pro: CCA, Gln: CAA, Arg: AGA,

Ser: TCT or TCC, Thr: ACT or ACC, Val: GTT or GTC,

Trp: TGG, Tyr: TAC

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As the foreign protein in this invention, human serum albumin, interferon- α , - β , or - γ , urokinase, growth hormone, insulin, Factor VIII, EPO, h-ANP, M-CSF and various lymphokines may be used.

In the case of human serum albumin, pre type, pro type, or prepro type may be used, and in the case of urokinase, pro type or any other type may be used. Among foreign proteins, in particular, a mature human serum albumin gene is preferable. According to the present invention, in the case that the mature human serum albumin gene is positioned immediately downstream to the serum albumin signal peptide gene, a substantial quantity of albumin can be produced.

Such foreign protein genes have been described in Japanese Patent Application (OPI) No. 29985/87 or EP-A-206733 (human serum albumin), Japanese Patent Application (OPI) No. 185189/86 or DE-A-3603958 (interferon- α), Japanese Patent Application (OPI) No. 108397/86 or EP-A-190686 (interferon- γ), Japanese Patent Application (OPI) No. 180591/85 or EP-A-154272 (urokinase), EP-A-160457 (Factor VIII), EP-A-148605 (EPO), WO85-4670 (h-ANP), WO86-4607 (M-CSF), and others.

In the above publications, the inventions are described as plasmids containing foreign protein genes.

The recombinant DNA for transforming yeast in this invention is prepared by linking the foreign protein gene downstream to the serum albumin signal peptide gene.

The promoter and terminator are not specifically limited to those found in yeast.

Acceptable promoters include PGK promoter (Nucleic Acid Res., 10(23), 7791 (1982)), ADH promoter (ibid.), *phoE* (5) promoter (J. Mol. Biol., 163(4), 513 (1983)), GAL1 promoter (Mol. Cell. Biol., 4(11), 2467 (1984)), GAL10 promoter (EP-A-132309) and GAP-DH promoter (J. Biol. Chem., 258, 5291 (1983)). Among these promoters, GAL1 promoter is particularly preferable.

The promoter is positioned upstream to the serum albumin signal peptide gene.

Acceptable terminators include the *phoE*(5) terminator (Cell, 12, 721-732 (1977)) and the GAP-DH terminator (J. Biol. Chem., 254, 9839-9845 (1979)).

The terminator is positioned downstream to the foreign protein gene.

The promoter and terminator may be obtained in a form already incorporated into plasmids.

The plasmid DNA must be capable of self-replication in yeast.

Acceptable examples are pJDB207 (Amersham) and pJDB219 (Amersham).

The recombinant plasmid of this invention is obtained either by cleaving a DNA sequence composed of the serum albumin signal peptide gene-foreign protein gene, a DNA sequence containing the promoter, and a DNA sequence containing the terminator from the above plasmid groups by a restriction enzyme and coupling (connecting) them to incorporate them into a proper plasmid, or by cleaving one DNA sequence and then incorporating it into another plasmid.

Also, the recombinant chromosome of this invention is obtained by insertion of a DNA sequence comprising the serum albumin signal peptide gene-foreign protein gene, a DNA sequence containing the promoter, and a DNA sequence containing the terminator into the yeast chromosome. The detail methods have been described in Proc. Natl. Acad. Sci. USA, 78, 6354-6358 (1981) and Method Enzymol., 101, 228-245 (1983).

The DNA sequence on the plasmid or the chromosome is arranged, from upstream to downstream, in the order of the promoter, serum albumin signal peptide gene, foreign protein gene, and terminator.

As the marker for selecting the desired plasmid, it is also possible to incorporate an antibiotic (tetracycline, ampicillin, kanamycin) resistance gene, or a gene to compensate for a nutritional requirement of the host. The method of preparing a transformant by this recombinant plasmid or the method of preparing a foreign protein is as follows.

The recombinant plasmid is introduced into the host cell i.e., yeast. Practically, a strain having a variation which is complemented by the selective marker gene carried by the plasmid to be inserted, for example, *Saccharomyces cerevisiae* AH22 (a, his4, leu2, can1) which is a leucine-requiring variant is acceptable for use.

Transformation of the host cell (yeast) is conducted by an established method, for example, the calcium phosphate sedimentation method, protoplast-polyethylene glycol fusion method, electroporation method.

The transformant is incubated in an established culture medium for the growth of the host cell. Practical examples of culture medium are YNB liquid culture medium (0.7 w/v% yeast nitrogen base (Difco Co.) and 2 w/v% glucose), YPD liquid culture medium (1 w/v% yeast extract (Difco), 2 w/v% polypeptone (Daigo Eiyo Sha), 2 w/v% glucose) and others.

Incubation is performed for 20 to 100 hours, usually at 15 to 43 °C (preferably about 30 °C), while being aerated or stirred as required.

After cultivation, the culture supernatant is recovered, and the foreign protein is purified by an established method, such as affinity chromatography or fractionation.

By using the method of this invention, a desired foreign protein can be produced by secretory expression. Compared with the system intracellular accumulation, production of the protein possessing structure and function much close to the native protein may be expected.

Additionally, in the system of intracellular accumulation, it is necessary to destroy the cells to refine the protein and to purify the protein from the liquid which contains debris, but this type of purification process is unnecessary when the method of this invention is used.

The use of the serum albumin signal peptide in expression of the protein also allows the development of the new secretory expression method. This increases the potential usefulness of this invention consider-

ably.

This invention is described in further detail below by referring to the following Example, which, however, is not intended to limit this invention in any respect.

Many of the techniques, reactions and analytical methods used in this invention are well known in the art. Unless otherwise specified, all enzymes can be obtained from commercial supply sources: for example, Takara Shuzo, Japan; New England Biolabs (NEB), Massachusetts, USA; Amersham, England; and Bethesda Research Laboratories (BRL), Maryland, USA.

Buffer solutions for enzymatic reactions and reaction conditions conformed to the recommended specifications of the manufacturers of the enzymes unless otherwise noted.

The transformation method of *Escherichia coli* by plasmid, colony hybridization, electrophoresis, and DNA recovery method from gels were conducted in accordance with the methods mentioned in "Molecular Cloning", Cold Spring Harbor Laboratory (1982). Yeast was transformed by the method stated in "Method in Yeast Genetics", Cold Spring Harbor Laboratory (1981).

EXAMPLE

Cloning of yeast GAL1, 10 promoters

(A) Preparation of yeast chromosomal DNA library

The chromosomal DNA of the yeast *Saccharomyces cerevisiae* GRF18 PHO80 *cir*⁰ strain (as described in EP-A-0180958) was extracted and purified by the method described by R. Cryer et al., (Method Enzymol., 12, 39 (1975)).

According to M. Mohnson and R.W. Davis (Mol. Cell. Biol., 4, 1440-1448 (1984)), the yeast GAL1, 10 promoter regions are located on the yeast chromosome, and when it is digested by the restriction enzymes *EcoRI* and *XbaI*, DNA segments of about 1 kb are obtained. Hence, yeast chromosomal DNA, extracted and purified as described above, was digested by *EcoRI* and *XbaI*, and DNA segments of about 1 kb were isolated by electrophoresis. These segments were mixed with plasmid pUC19 (BRL) which was digested by *EcoRI* and *XbaI*, and dephosphorylated at its 5' terminal with alkaline phosphatase derived from calf intestines (CIP). These were ligated using the ligation kit (Takara Shuzo). This product was introduced into *Escherichia coli* JM109 (Takara Shuzo). The transformant was applied to a YT agar plate containing 0.004 w/v% X-gal (5-bromo-4-chloro-3-indolyl- β -galactoside) and 1 mM IPTG (isopropyl- β ,D-thiogalactopyranoside), and was incubated overnight at 37 °C. (To prepare the agar plate 8 g of polypeptone, 5 g of yeast extract, and 5 g of sodium chloride were dissolved in water to make up 1 liter and 12 g of agar powder was added. After sterilization in an autoclave, the mixture was dispensed into plastic Petri dish and solidified; X-gal and IPTG were added after autoclaving once the culture medium had cooled.)

White and blue colonies appeared, and only the white colonies having the DNA inserts were used. (The desired transformant produced white colonies since the recombinant plasmid inserted therein had no lac Z gene.) One hundred colonies were inoculated onto an L-agar plate containing 40 μ g/ml ampicillin by a sterilized toothpick. (To prepare the agar plate 0.62 g of tris base, 10 g of polypeptone, 5 g of yeast extract, and 5 g of sodium chloride were dissolved in water to make up 1 liter, and 12 g of agar powder was added. The mixture was sterilized in an autoclave, dispensed into plastic Petri dish and solidified; ampicillin added, after autoclaving once the medium had cooled.) This L-agar plate was incubated overnight at 37 °C. By this method, a library consisting of about 5,000 colonies was prepared. The formed colonies were transferred to a nitrocellulose filter, dipped in a solution of 0.5 M sodium hydroxide and 1.5 M sodium chloride to denature the DNA, and were neutralized in a solution of 1.5 M sodium chloride and 0.5 M tris-hydrochloric acid at pH 7.5. The *E. coli* debris was washed with 2 x SSC (0.3 M sodium chloride, 0.03 M sodium citrate at pH 7.0) and removed, and after drying the filter in air, it was subjected to vacuum drying for 2 hours at 80 °C.

(B) Preparation of the probe

Part of base sequence of the gene coding for the GAL1, 10 promoters was synthesized by the phosphoramidite method using a DNA synthesizer, Applied Biosystem Co. model 381A. Its sequence is shown below.

5'-CTCTATACTTTAACGTCAAG-3'

The sequence was subjected to electrophoresis using 7 M urea-20 w/v% polyacrylamide gel and purified. The 5' terminal of the purified DNA sequence was labeled radioactively by [γ -³²P] ATP and T4 polynucleotide kinase. The reaction using 10 pmoles of synthetic DNA, 250 μ Ci of [γ -³²P] ATP, and 8 units

of T4 polynucleotide kinase, resulted in a synthetic DNA probe terminally labeled with ^{32}P (2×10^7 cpm (Cerenkov count)). The synthetic DNA probe was purified by NENSORB 20 (Du Pont).

(C) Screening of GAL1, 10 promoters

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Nitrocellulose filters having the DNA fixed as described in step (A) were placed in vinyl bags with each set containing 10 filters, and the following process carried out. Ten milliliters of prehybridization solution composed of 6 x SSC, 0.1 w/v% SDS, and 20 $\mu\text{g}/\text{mL}$ of salmon sperm DNA cooled on ice after heating for 5 minutes at 100°C was put in a vinyl bag which was sealed and incubated for 3 hours at 40°C . The prehybridization solution was then discarded and 10 mL of hybridization solution was added and incubated overnight at 40°C . The hybridization solution contained 6 x SSC, 0.1 w/v% SDS, 100 $\mu\text{g}/\text{mL}$ salmon sperm DNA, and 7.5×10^5 cpm/mL ^{32}P -probe. After incubation, the filter was transferred to a beaker and washed in 6 x SSC, 0.1 w/v% SDS for 30 minutes at 50°C , in 2 x SSC and 0.1 w/v% SDS for 30 minutes at 50°C , in 2 x SSC and 0.1 w/v% SDS for 30 minutes at 50°C , and finally in 0.1 x SSC and 0.1 w/v% SDS for 30 minutes at 50°C . The washed filter was dried in air and subjected to autoradiography after applying spotting marks of 100-200 cpm. As a result, two positive clones were obtained. One of the clones was subjected to shaking culture overnight at 37°C in super broth containing 40 $\mu\text{g}/\text{mL}$ of ampicillin. (To prepare the super broth 12 g of bactotrypton, 24 g of yeast extract, and 5 mL of glycerol were dissolved in water to make up 900 mL, which was sterilized by autoclave to obtain solution A. Then, 3.81 g of potassium dihydrogen phosphate and 12.5 g of potassium monohydrogen phosphate were dissolved in water to make up 100 mL, which was sterilized by autoclave to obtain solution B. These solutions A and B were mixed in a ratio of 9:1 by v/v.) Then, the plasmid DNA was extracted and purified by the alkaline-SDS method.

When part of the base sequence of this plasmid DNA (pGAL11, Fig. 1) was examined by the dideoxy method, the results coincided with the reported sequence by M. Johnston and R.W. Davis (Mol. Cell, Biol., 4, 1440-1448, (1984)). That is, it was found that pGAL11 possessed the GAL1 promoter in the direction of the XbaI site from the EcoRI site, and the GAL10 promoter in the opposite direction.

(D) Conversion of pGAL11 from the XbaI site to the BamHI site

When ligating the promoter sequence on pGAL11 with the DNA sequence coding for the signal peptide and human serum albumin, it is not convenient to have an intervening XbaI site because the XbaI site is present on the human serum albumin gene. Therefore, the XbaI site was converted to the BamHI site as follows.

After digesting pGAL11 by XbaI, the sticky end was repaired by *E. coli*-derived DNA polymerase I, Klenow fragment, in the presence of dGTP, dATP, dTTP, dCTP. To this DNA fragment, the BamHI linker pCGGATCCG having a phosphorylated 5' terminal was added and was ligated by T4 DNA ligase. After then digesting with BamHI, ligation was again carried out with T4 DNA ligase and the resulting plasmid introduced into *E. coli* HB101 (EP-A-13828). From the resulting transformants, a clone having plasmid pGAL12 (as shown in Fig. 1) was obtained. By digesting pGAL12 by EcoRI and BamHI, the GAL1 and GAL10 promoters could be isolated as a DNA fragment of about 1 kb.

(E) Preparation of *E. coli*-yeast shuttle vector pPT2 possessing a yeast pho5 terminator

The plasmid pAP5 which has encoded the *Saccharomyces cerevisiae* pho5 gene is disclosed in Japanese Patent Application (OPI) No. 151183/87 or EP-A-216573. This plasmid was digested by the restriction enzymes Sau3AI and PstI, and the DNA fragment which has encoded the pho5 terminator, about 370 bp, was isolated by electrophoresis (Fig. 2). The commercially available pUC9 (BRL) was then digested with BamHI and PstI, treated with alkaline phosphatase, and ligated with the 370 bp DNA fragment. The base sequence at the Sau3AI cleavage site of the 370 bp fragment was

GATCC.....
G.....

and when it was ligated with the sticky end of the BamHI, the BamHI site was regenerated. Therefore, by digesting plasmid pPT1 obtained in the above ligation reaction with BamHI and PstI, or by digestion with BamHI and HindIII, a DNA fragment possessing a 370 bp pho5 terminator was obtained (Fig. 2).

The commercially available shuttle vector pJDB207 (Fig. 3) is self-replicating in *E. coli* and yeast. After digestion with BamHI and HindIII, it was treated with alkaline phosphatase. After digesting pPT1 with BamHI and HindIII, the DNA fragment having the 370 bp pho5 terminator was isolated by electrophoresis and was ligated with pJDB207. From the resulting transformants, a clone having plasmid pPT2 (as shown in Fig. 3)

was obtained. pPT2 is an *E. coli*-yeast shuttle vector possessing a *pho5* terminator. In *E. coli*, it possesses an ampicillin resistance marker with β -lactamase activity and in yeast it has a marker to compensate for a leucine nutritional requirement.

5 (F) Human serum albumin gene

The DNA sequence coding for human serum albumin was derived from the plasmid pGX401 (Figs. 4 and 5) disclosed in Japanese Patent Application (OPI) No. 29985/87 or EP-A-206733 as follows. pGX401 was digested with the restriction enzymes *Xba*I and *Hind*III, and the DNA fragment (HSA2) of about 750 bp
10 coding for the C-terminal side ³⁵⁷Leu to ⁵⁸⁵Leu of the amino acid sequence of human serum albumin, including the 3' untranslated region, was isolated by electrophoresis. The commercially available plasmid pUC19 was digested with *Xba*I and *Hind*III, was treated with alkaline phosphatase to dephosphorylate the 5' terminal and was ligated with HSA2 with T4 DNA ligase. It was introduced into *E. coli* HB101, and from the resulting transformants, a clone having plasmid pHSA2 (as shown in Fig. 6) was obtained.

15 Upon digesting pGX401 with *Dra*I and *Xba*I, a DNA fragment of about 1 kb was isolated by electrophoresis. This DNA fragment is the DNA sequence encoding for the N-terminal side ¹²Lys to ³⁵⁶Thr of the amino acid sequence of human serum albumin.

Using the DNA synthesizer Applied Biosystem model 381A, the following DNA sequence encoding for the N-terminal ¹Asp to ¹¹Phe of the amino acid sequence of mature human serum albumin was synthesized
20 by the phosphoramidite method.

	1	2	3	4	5	6	7	8	9	10	11
	Asp	Ala	His	Lys	Ser	Glu	Val	Ala	His	Arg	Phe
25 TC	GAC G	GCA CGT	CAC GTG	AAG TTC	AGT TCA	GAG CTC	GTT CAA	GCT CGA	CAT GTA	CGG GCC	TTT AAA

The codon for aspartic acid (Asp) in pGX401 was GAT, but GAC was used here. As a result, after
30 ligating the synthetic DNA with the 1 kb DNA fragment derived from pGX401, the *Sal*I site was regenerated when it was inserted into the *Sal*I-*Xba*I site of pUC19. Furthermore, when digested with *Hin*CII, the DNA sequence coding for the amino acid sequence starting from the N-terminal ¹Asp of mature human serum albumin was obtained.

The 5' terminal of the synthetic DNA was phosphorylated by ATP and T4 polynucleotide kinase.
35 pGX401 was digested with *Dra*I and *Xba*I and 1 kb DNA fragment was isolated by electrophoresis. This fragment and the phosphorylated synthetic DNA were ligated with T4 ligase, digested with *Sal*I and *Xba*I, and then ligated with pUC19 which was digested with *Sal*I and *Xba*I and dephosphorylated by CIP. The resulting DNA was introduced into *E. coli* HB101, and from the transformants, a clone having the plasmid pHSA1 (as shown in Fig. 7) was obtained.

40

(G) Preparation of plasmid DNA for expressing and secreting human serum albumin in yeast

The DNA sequence shown in Table 3 below coding for the signal peptide of human serum albumin was synthesized by the phosphoramidite method by the DNA synthesizer Applied Biosystem model 381A.

45

50

55

Table 3

Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe Ser Scr Ala Tyr Scr

GATCCACA ATG AAA TGG GTT ACT TTC ATT TCT TTG TTC TTG TCT TCT TCT GCT TAC TCT
GTGT TAC TTT ACC CAA TGA AAG TAA AGA AAC AAC AAG AAG AAG AAG CGA ATG AGA

Also, the DNA sequence encoding the signal peptide amino acid which was changed to Arg or His, Ala or Pro, Lys or Gly, Val or Cys, Trp or Ser, Ala or Gly in the place of -17, -5, -4, -3, -2 and -1, respectively, was synthesized by the same method (cf. Table 2). The changed DNA sequence lead to produce and
55 secrete the more proper N-terminal side of albumin.

The 5' terminal of the synthetic DNA was phosphorylated with ATP and T4 polynucleotide kinase. pHSA1 was digested with *Xba*I and *Hinc*II, and the 1kb HSA1 DNA fragment encoding for the N-terminal side of human serum albumin was isolated by electrophoresis. The phosphorylated synthetic DNA and

HSA1 were mixed and ligated with T4 DNA ligase, and digested further with XbaI and BamHI. After digesting pHSA2 with XbaI and BamHI, it was treated with alkaline phosphatase. After mixing, these DNAs were ligated with T4 DNA ligase and introduced into *E. coli* HB101 cells. Among the resulting transformants a clone having the plasmid pNH001 (as shown in Fig. 8) was obtained.

5 After digesting pNH001 with EcoRI and BamHI, it was treated with alkaline phosphatase. Then, pGAL12 was digested with EcoRI and BamHI, a DNA fragment of 1 kb possessing the GAL1 promoter was isolated by electrophoresis, mixed with the treated pNH001 and ligated with T4 DNA ligase. From the resulting transformants, a clone having the plasmid pNH007 (as shown in Fig. 9) was obtained. pNH007 is a plasmid DNA having the DNA sequence encoding for the human serum albumin signal peptide located downstream
10 from the GAL1 promoter, the DNA sequence encoding for mature human serum albumin immediately after it, and immediately following that, the 3' untranslated region derived from human serum albumin cDNA inserted in the EcoRI-HindIII site of pUC19.

After digesting pNH007 with EcoRI and HindIII, a DNA fragment of 2.7 kb coding for the GAL1 promoter, the signal peptide, mature human serum albumin and the untranslated region was isolated by
15 electrophoresis. Additionally, pPT2 was digested with BamHI and treated with alkaline phosphatase. It was mixed with the 2.7 kb DNA fragment, and the sticky end was repaired by DNA polymerase I, Klenow fragment, in the presence of dATP, dGTP, dTTP, and dCTP. After ligation with T4 DNA ligase, it was introduced into *E. coli* HB101. From the resulting transformants, a clone having the plasmid pNH008 (as shown in Fig. 10) was obtained.

20 pNH008 is a plasmid capable of self-replication in *E. coli* and yeast and possesses the DNA sequence encoding for the human serum albumin signal peptide and the succeeding mature human serum albumin protein under the control of the GAL1 promoter functionable in yeast. Furthermore, pNH008 also possesses a gene for ampicillin resistance in *E. coli*, and a gene for fulfilling the nutritional requirement for leucine in yeast, and these genes can be used as selective marker for transformants.

25 (H) Introduction of plasmid pNH008 into Yeast

Plasmid pNH008, for the secretory expression of human serum albumin, was introduced into yeast *Saccharomyces cerevisiae* AH22 (Proc. Natl. Acad. Sci. USA, 75, 1929-1933 (1978)) by the following
30 method.

S. cerevisiae AH22 was subjected to shaking culture overnight at 30 °C in 50 ml of YPD medium. (To prepare the medium, 10 g of yeast extract and 20 g of bactopectone were dissolved in water to make up 900 ml, which was sterilized in an autoclave and mixed with 100 ml of 20 w/v% glucose separately
35 sterilized in an autoclave). The cells were precipitated by centrifugation, resuspended in 20 ml of water, and centrifuged again. Next, the cells were suspended in 10 ml of 50 mM dithiothreitol, 1.2 M sorbitol, 2 mM EDTA at pH 8.5, and were shaken slowly for 10 minutes at 30 °C. The cells were collected by centrifugation, and suspended in 10 ml of 1.2 M sorbitol, then centrifuged again for collection. The cells were suspended in 10 ml of 0.2 mg/ml zymolyase 100T, 1.2 M sorbitol, 10 mM EDTA, 0.1 M sodium citrate at pH 5.8, and were shaken slowly for 1 hour at 30 °C. The cells were collected by centrifugation and
40 washed in 10 ml each of 1.2 M sorbitol, 10 mM calcium chloride and 1.2 M sorbitol, sequentially, and again the cells were collected by centrifugation. The cells were suspended in 1 ml of 10 mM calcium chloride and 1.2 M sorbitol. One hundred microliter aliquotes of suspension were placed in a sterile test tube and mixed with 5 µl (5 µg) of pNH008; the mixture was allowed to stand for 15 minutes at room temperature. After this, it was mixed with 1.2 ml of 20 w/v% polyethylene glycol 4,000, 10 mM calcium chloride, 10 mM tris-hydrochloride at pH 7.5, and after gentle mixing, the mixture was let stand at room
45 temperature for 20 minutes. The cells were collected by centrifugation, suspended in 0.1 ml of YPD medium containing 1.2 M sorbitol and 10 mM calcium chloride, and shaken gently for 30 minutes at 30 °C. 1, 5, 10, 20 and 50 µl of suspension were suspended in 45 °C-controlled 10 ml of 1.2 M sorbitol, 3 w/v% noble agar, 2 w/v% glucose, and 0.7 w/v% yeast nitrogen base and were spread over plates composed of
50 1.2 M sorbitol, 3 w/v% bactoagar, 2 w/v% glucose, and 0.7 w/v% yeast nitrogen base. After the plates solidified, they were subjected to stationary culture for 3 days at 30 °C. Formed colonies were collected by a sterile toothpick suspended in 3 ml of 0.7 w/v% yeast nitrogen base and 2 w/v% glucose, and subjected to shaking culture for 2 days at 30 °C. One and a half milliliters of suspension was centrifuged, and the cells were collected and suspended in 3 ml of YPG medium. (To prepare the culture, 10 g of yeast extract and
55 20 g of bactopectone were dissolved in water to make up 900 ml, sterilized in an autoclave, and mixed with 100 ml of 20 w/v% galactose, sterilized separately in an autoclave.) This was subjected to shaking culture at 30 °C. The human serum albumin concentration in the culture supernatant was measured by the RPHA method (as described in European Patent 122,620), and a maximum human serum albumin of 10

μg/ml was detected on the first day.

(I) Cultivation of yeast for the expression and secretion of human serum albumin

- 5 The yeast *S. cerevisiae* AH22 for the expression and secretion human serum albumin transformed by pNH008 as mentioned above was cultivated by the following procedure. The recombinant yeast was grown in a plate containing 0.7 w/v% yeast nitrogen base, 2 w/v% glucose and 3 w/v% bactoagar and collected by a platinum loop. It was inoculated into YNB medium 50 ml composed of 0.7 w/v% yeast nitrogen base and 2 w/v% glucose and incubated for 2 days at 30 °C. The whole volume was inoculated into 500 ml of YNB
10 medium and incubated for 2 days at 30 °C. The cells were collected by centrifugation, and suspended in 500 ml of YPG medium, and subjected to shaking culture at 30 °C. A portion of the culture broth was collected after 0, 3, 6, 24 and 48 hours of incubation, and the culture supernatant was obtained by centrifugation. The concentration of human serum albumin secreted into the culture broth was measured by the RPHA method. Secretory expression of human serum albumin was detected beginning the third hour
15 after the start of incubation, and the concentration of human serum albumin in the supernatant was 0.25 mg/l at 6 hours, 20 mg/l at 24 hours, and 160 mg/l at 48 hours.

Claims

Claims for the following Contracting States : BE, CH, DE, FR, GB, IT, LI, NL, SE

20

1. A serum albumin signal peptide gene encoding the following amino acid sequence:

25

Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe
X₅ X₄ X₃ X₂ X₁

30

wherein Y represents Lys, Arg or His; X₅ represents Ala, Pro or Ser; X₄ represents Lys, Gly or Ser; X₃ represents Ala, Val or Cys; X₂ represents Tyr, Trp or Ser; and X₁ represents Ser, Ala or Gly;

with the proviso that the gene does not encode the amino acid sequence

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerAlaTyrSer

35

2. A method for preparing a foreign protein comprising expressing and secreting said foreign protein by yeast transformed by a recombinant DNA comprising the serum albumin signal peptide gene according to claim 1 adjacent to the gene of said foreign protein, a promoter upstream to said serum albumin signal peptide gene and a terminator downstream to said foreign protein gene.

40

3. A recombinant DNA for transforming yeast comprising the serum albumin signal peptide gene according to claim 1 adjacent to DNA encoding a foreign protein, a promoter upstream to the serum albumin signal peptide gene and a terminator downstream to the foreign protein gene.

4. A strain of yeast transformed by a recombinant DNA according to claim 3.

45

5. A method according to claim 2, wherein the amino acid sequence of said serum albumin signal peptide is selected from the group consisting of:

50

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerValTyrAla,

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheAlaLysValSerAla,

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheProGlyCysTrpAla,

and

55

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheProGlyValTrpAla.

6. A method according to claim 2, wherein said foreign protein is selected from the group consisting of human serum albumin, interferon- α , interferon- β , interferon- γ , urokinase, growth hormone, insulin, lymphokines, h-ANP, Factor VIII, CSFs and EPO.

5 7. A method according to claim 2, wherein the foreign protein gene is the mature human serum albumin gene which is positioned immediately downstream to the serum albumin signal peptide gene.

Claims for the following Contracting State : ES

10 1. A method for preparing a foreign protein comprising expressing and secreting said foreign protein by yeast transformed by a recombinant DNA comprising the serum albumin signal peptide gene adjacent to the gene of said foreign protein, a promoter upstream to said serum albumin signal peptide gene and a terminator downstream to said foreign protein gene;
wherein the serum albumin signal peptide gene encodes the following amino acid sequence:

Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe
X₅ X₄ X₃ X₂ X₁

20 wherein Y represents Lys, Arg or His; X₅ represents Ala, Pro or Ser; X₄ represents Lys, Gly or Ser; X₃ represents Ala, Val or Cys; X₂ represents Tyr, Trp or Ser; and X₁ represents Ser, Ala or Gly;
with the proviso that the gene does not encode the amino acid sequence
MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerAlaTyrSer

25 2. A method according to claim 1, wherein the amino acid sequence of said serum albumin signal peptide is selected from the group consisting of:

30 MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerValTyrAla,

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheAlaLysValSerAla,

35 MetLysTrpValThrPheIleSerLeuLeuPheLeuPheProGlyCysTrpAla,

and

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheProGlyValTrpAla.

40 3. A method according to claim 1, wherein said foreign protein is selected from the group consisting of human serum albumin, interferon- α , interferon- β , interferon- γ , urokinase, growth hormone, insulin, lymphokines, h-ANP, Factor VIII, CSFs and EPO.

45 4. A method according to claim 1, wherein the foreign protein gene is the mature human serum albumin gene which is positioned immediately downstream to the serum albumin signal peptide gene.

Patentansprüche

50 Patentansprüche für folgende Vertragsstaaten : BE, CH, DE, FR, GB, IT, LI, NL, SE

1. Serumalbumin-Signalpeptid-Gen, das für folgende Aminosäuresequenz codiert:

55 Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe X₅ X₄ X₃ X₂ X₁

wobei Y für Lys, Arg oder His steht; X₅ für Ala, Pro oder Ser steht; X₄ für Lys, Gly oder Ser steht; X₃ für Ala, Val oder Cys steht; X₂ für Tyr, Trp oder Ser steht; und X₁ für Ser, Ala oder Gly steht; mit der Maßgabe, daß das Gen nicht für die folgende Aminosäuresequenz codiert:

5

**Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Ser Ser Ala Tyr Ser.**

10

2. Verfahren zur Herstellung eines fremden Proteins, umfassend die Expression und Absonderung des fremden Proteins durch Hefe, die durch eine rekombinante DNA transformiert ist, die das Serumalbumin-Signalpeptid-Gen nach Anspruch 1 benachbart zu dem Gen für das fremde Protein, einen Promotor in 5'-Richtung von dem Serumalbumin-Signalpeptid-Gen und einen Terminator in 3'-Richtung von dem Gen für das fremde Protein umfaßt.
3. Rekombinante DNA zur Transformation von Hefe, umfassend das Serumalbumin-Signalpeptid-Gen nach Anspruch 1 benachbart zu DNA, die für ein fremdes Protein codiert, einen Promotor in 5'-Richtung von dem Serumalbumin-Signalpeptid-Gen und einen Terminator in 3'-Richtung von dem Gen für das fremde Protein.
4. Hefestamm, der mit der rekombinanten DNA nach Anspruch 3 transformiert ist.
5. Verfahren nach Anspruch 2, wobei die Aminosäuresequenz des Serumalbumin-Signalpeptids aus der folgenden Gruppe ausgewählt ist:

30

**Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Ser Ser Val Tyr Ala**

35

**Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Ala Lys Val Ser Ala**

40

und

45

**Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Pro Gly Val Trp Ala.**

50

6. Verfahren nach Anspruch 2, wobei das fremde Protein unter menschlichem Serumalbumin, Interferon- α , Interferon- β , Interferon-gamma, Urokinase, Wachstumshormon, Insulin, Lymphokinen, h-ANP, Faktor VIII, CSFs und EPO ausgewählt ist.
7. Verfahren nach Anspruch 2, wobei es sich bei dem Gen für das fremde Protein um das Gen für reifes menschliches Serumalbumin handelt, das unmittelbar in 3'-Position von dem Serumalbumin-Signalpeptid-Gen angeordnet ist.

55

Patentansprüche für folgenden Vertragsstaat : ES

1. Verfahren zur Herstellung eines fremden Proteins, umfassend die Expression und Absonderung des fremden Proteins durch Hefe, die durch eine rekombinante DNA transformiert ist, die das Serumalbumin-Signalpeptid-Gen benachbart zu dem Gen für das fremde Protein, einen Promotor in 5'-Richtung von dem Serumalbumin-Signalpeptid-Gen und einen Terminator in 3'-Richtung von dem Gen für das fremde Protein umfaßt;
wobei das Serumalbumin-Signalpeptid-Gen für folgende Aminosäuresequenz codiert:

Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe X₅ X₄ X₃ X₂ X₁

wobei Y für Lys, Arg oder His steht; X₅ für Ala, Pro oder Ser steht; X₄ für Lys, Gly oder Ser steht; X₃ für Ala, Val oder Cys steht; X₂ für Tyr, Trp oder Ser steht; und X₁ für Ser, Ala oder Gly steht; mit der Maßgabe, daß das Gen nicht für die folgende Aminosäuresequenz codiert:

Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Ser Ser Ala Tyr Ser.

2. Verfahren nach Anspruch 1, wobei die Aminosäuresequenz des Serumalbumin-Signalpeptids aus der folgenden Gruppe ausgewählt ist:

Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Ser Ser Val Tyr Ala

Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Ala Lys Val Ser Ala

Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Pro Gly Cys Trp Ala

und

Met Lys Trp Val Thr Phe Ile Ser Leu Leu Phe Leu
Phe Pro Gly Val Trp Ala.

3. Verfahren nach Anspruch 1, wobei das fremde Protein unter menschlichem Serumalbumin, Interferon- α , Interferon- β , Interferon-gamma, Urokinase, Wachstumshormon, Insulin, Lymphokinen, h-ANP, Faktor VIII, CSFs und EPO ausgewählt ist.
4. Verfahren nach Anspruch 1, wobei es sich bei dem Gen für das fremde Protein um das Gen für reifes menschliches Serumalbumin handelt, das unmittelbar in 3'-Position von dem Serumalbumin-Signalpeptid-Gen angeordnet ist.

Revendications

Revendications pour les Etats contractants suivants : BE, CH, DE, FR, GB, IT, LI, NL, SE

1. Gène du peptide signal de la sérumalbumine codant la séquence d'acides aminés suivante :

Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe
X₅ X₄ X₃ X₂ X₁

dans laquelle Y représente Lys, Arg ou His ; X₅ représente Ala, Pro ou Ser ; X₄ représente Lys, Gly ou Ser ; X₃ représente Ala, Val ou Cys ; X₂ représente Tyr, Trp ou Ser et X₁ représente Ser, Ala ou Gly ; à condition que le gène ne code pas la séquence d'acides aminés :

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerAlaTyrSer.

2. Procédé de préparation d'une protéine étrangère comprenant l'expression et la sécrétion de ladite protéine étrangère par une levure transformée par un ADN recombiné comprenant le gène du peptide signal de la sérumalbumine selon la revendication 1 en position adjacente au gène de ladite protéine étrangère, un promoteur en amont dudit gène du peptide signal de la sérumalbumine et un terminateur en aval dudit gène de la protéine étrangère.

3. ADN recombiné pour transformer une levure, comprenant le gène du peptide signal de la sérumalbumine selon la revendication 1 en position adjacente à un ADN codant une protéine étrangère, un promoteur en amont du gène du peptide signal de la sérumalbumine et un terminateur en aval du gène de la protéine étrangère.

4. Souche de levure transformée par un ADN recombiné selon la revendication 3.

5. Procédé selon la revendication 2, dans lequel la séquence d'acides aminés dudit peptide signal de la sérumalbumine est choisie dans le groupe formé par :

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheSerSerValTyrAla,

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheAlaLysValSerAla,

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheProGlyCysTrpAla,

et

MetLysTrpValThrPheIleSerLeuLeuPheLeuPheProGlyValTrpAla.

6. Procédé selon la revendication 2, dans lequel ladite protéine étrangère est choisie dans le groupe formé par la sérumalbumine humaine, l'interféron α , l'interféron β , l'interféron γ , l'urokinase, l'hormone de croissance, l'insuline, les lymphokines, h-ANP, le facteur VIII, les CSF et EPO.

7. Procédé selon la revendication 2, dans lequel le gène de la protéine étrangère est le gène de la sérumalbumine humaine mature qui est situé immédiatement en aval du gène du peptide signal de la sérumalbumine.

Revendications pour l'Etat contractant suivant : ES

1. Procédé de préparation d'une protéine étrangère comprenant l'expression à la sécrétion de ladite protéine étrangère par une levure transformée par un ADN recombiné comprenant le gène du peptide signal de la sérumalbumine en position adjacente au gène de ladite protéine étrangère, un promoteur en amont dudit gène du peptide signal de la sérumalbumine et un terminateur en aval dudit gène de la protéine étrangère, dans lequel le gène du peptide signal de la sérumalbumine code la séquence d'acides aminés suivante :

Met Y Trp Val Thr Phe Ile Ser Leu Leu Phe Leu Phe
X₅ X₄ X₃ X₂ X₁

5

dans laquelle Y représente Lys, Arg ou His ; X₅ représente Ala, Pro ou Ser ; X₄ représente Lys, Gly ou Ser ; X₃ représente Ala, Val ou Cys ; X₂ représente Tyr, Trp ou Ser et X₁ représente Ser, Ala ou Gly ; à condition que le gène ne code pas la séquence d'acides aminés :
MetLysTrpValThrPhelleSerLeuLeuPheLeuPheSerSerAlaTyrSer.

10

2. Procédé selon la revendication 1, dans lequel la séquence d'acides aminés dudit peptide signal de la sérumalbumine est choisie dans le groupe formé par :

15

MetLysTrpValThrPhelleSerLeuLeuPheLeuPheSerSerValTyrAla,
MetLysTrpValThrPhelleSerLeuLeuPheLeuPheAlaLysValSerAla,
MetLysTrpValThrPhelleSerLeuLeuPheLeuPheProGlyCysTrpAla,
et

20

MetLysTrpValThrPhelleSerLeuLeuPheLeuPheProGlyValTrpAla.

25

3. Procédé selon la revendication 1, dans lequel ladite protéine étrangère est choisie dans le groupe formé par la sérumalbumine humaine, l'interféron α , l'interféron β , l'interféron γ , l'urokinase, l'hormone de croissance, l'insuline, les lymphokines, h-ANP, le facteur VIII, les CSF et EPO.

30

4. Procédé selon la revendication 1, dans lequel le gène de la protéine étrangère est le gène de la sérumalbumine humaine mature qui est situé immédiatement en aval du gène du peptide signal de la sérumalbumine.

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Fig. 1

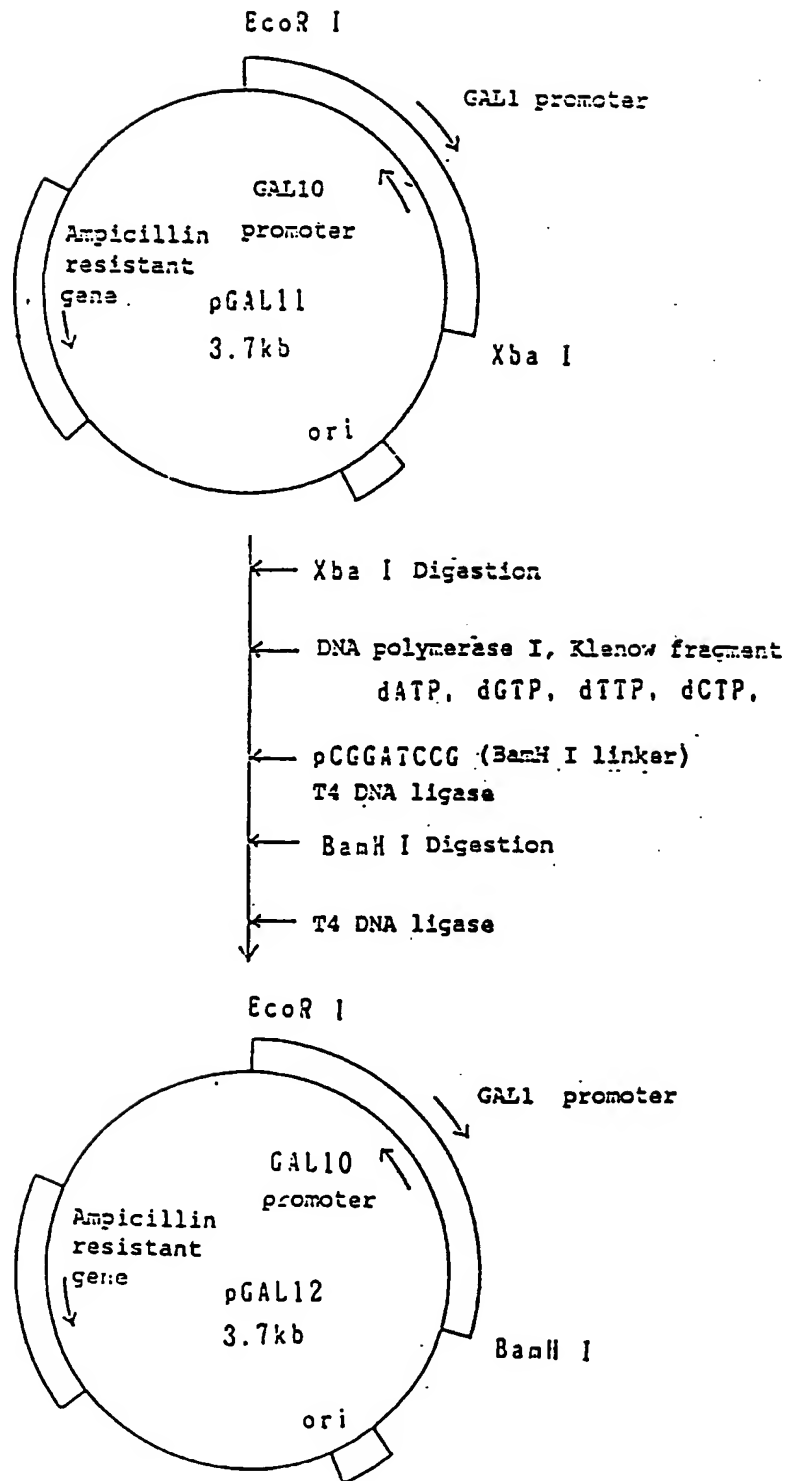


Fig. 2

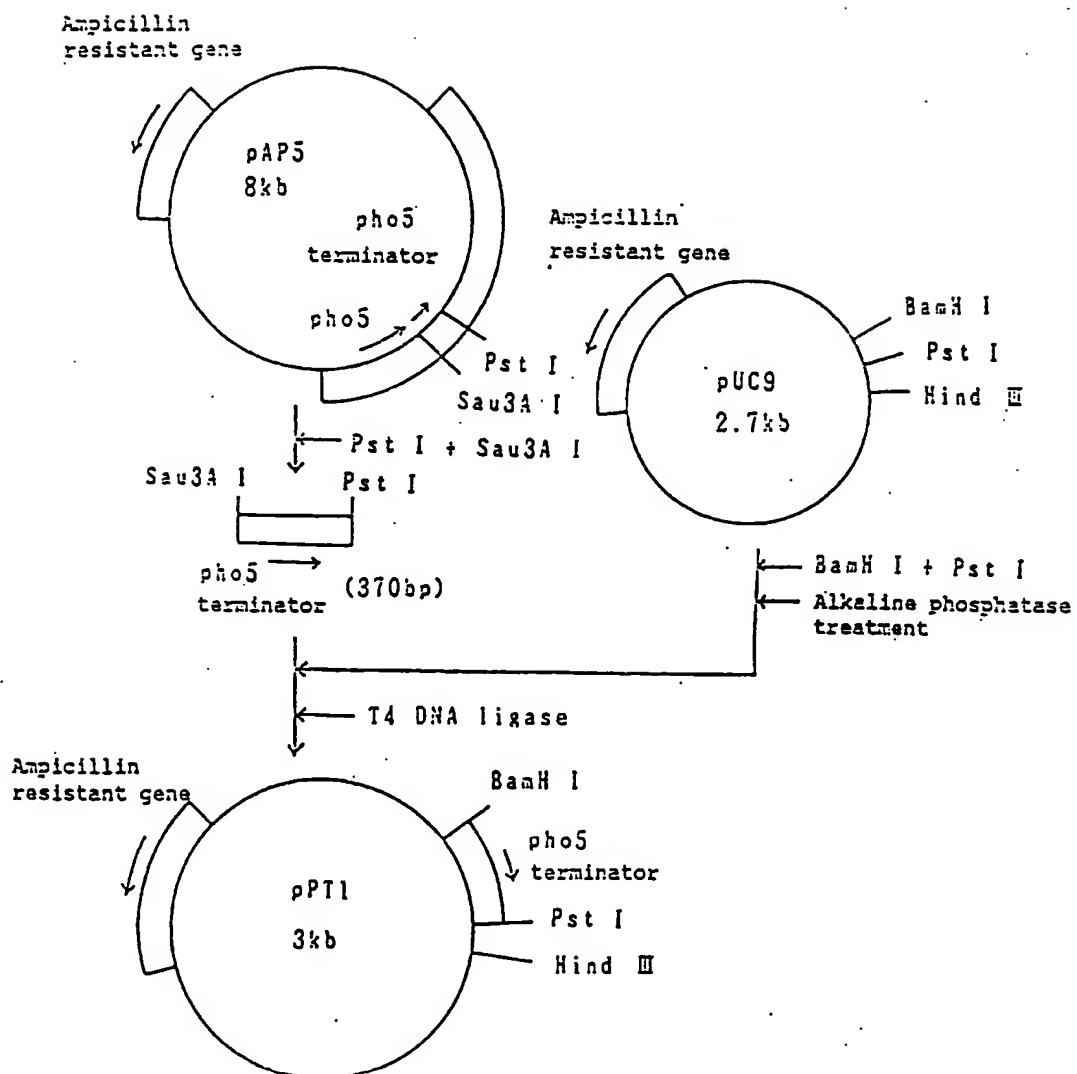


Fig. 3

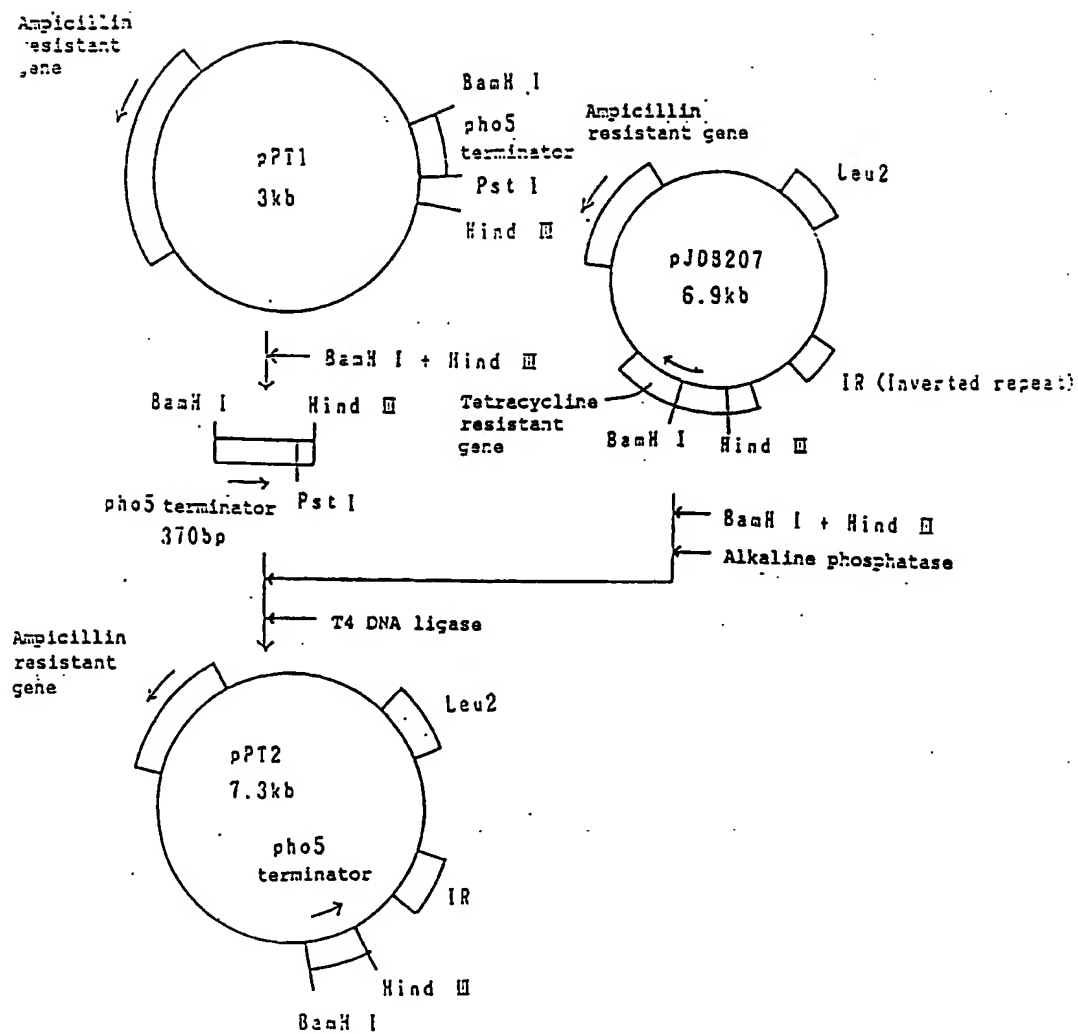


Fig. 4

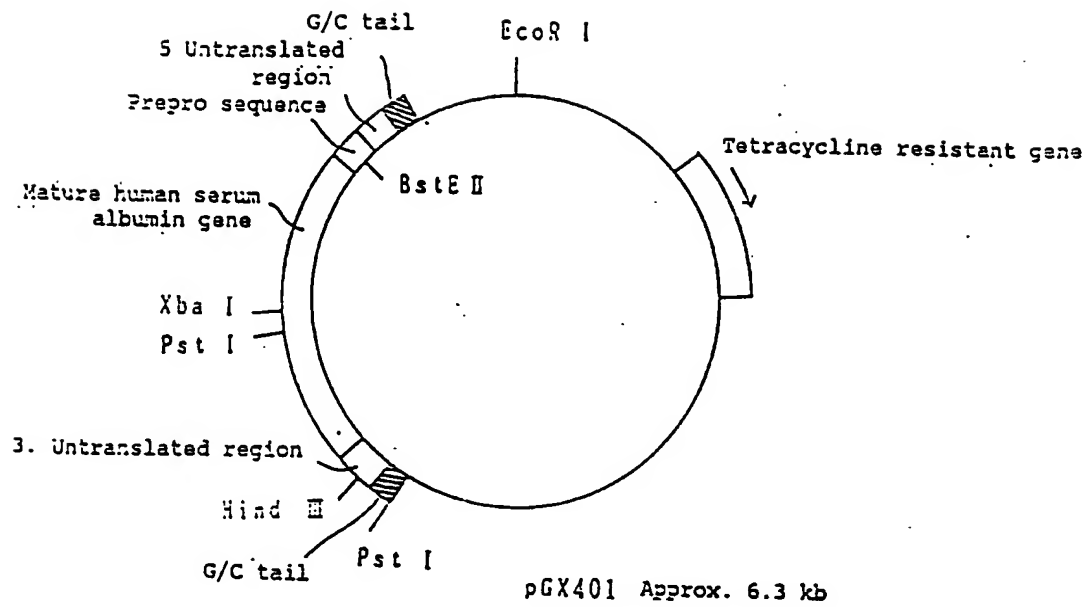


Fig. 5

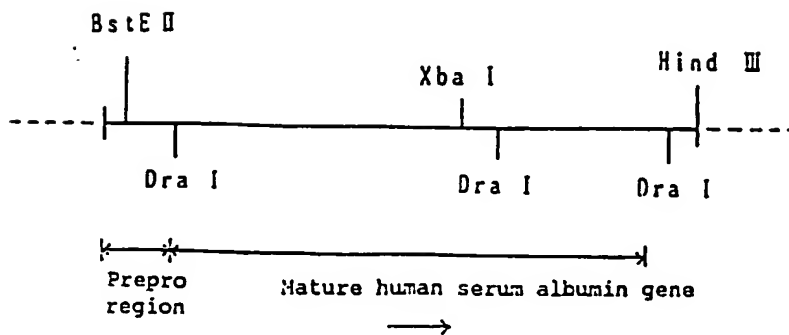


Fig. 6

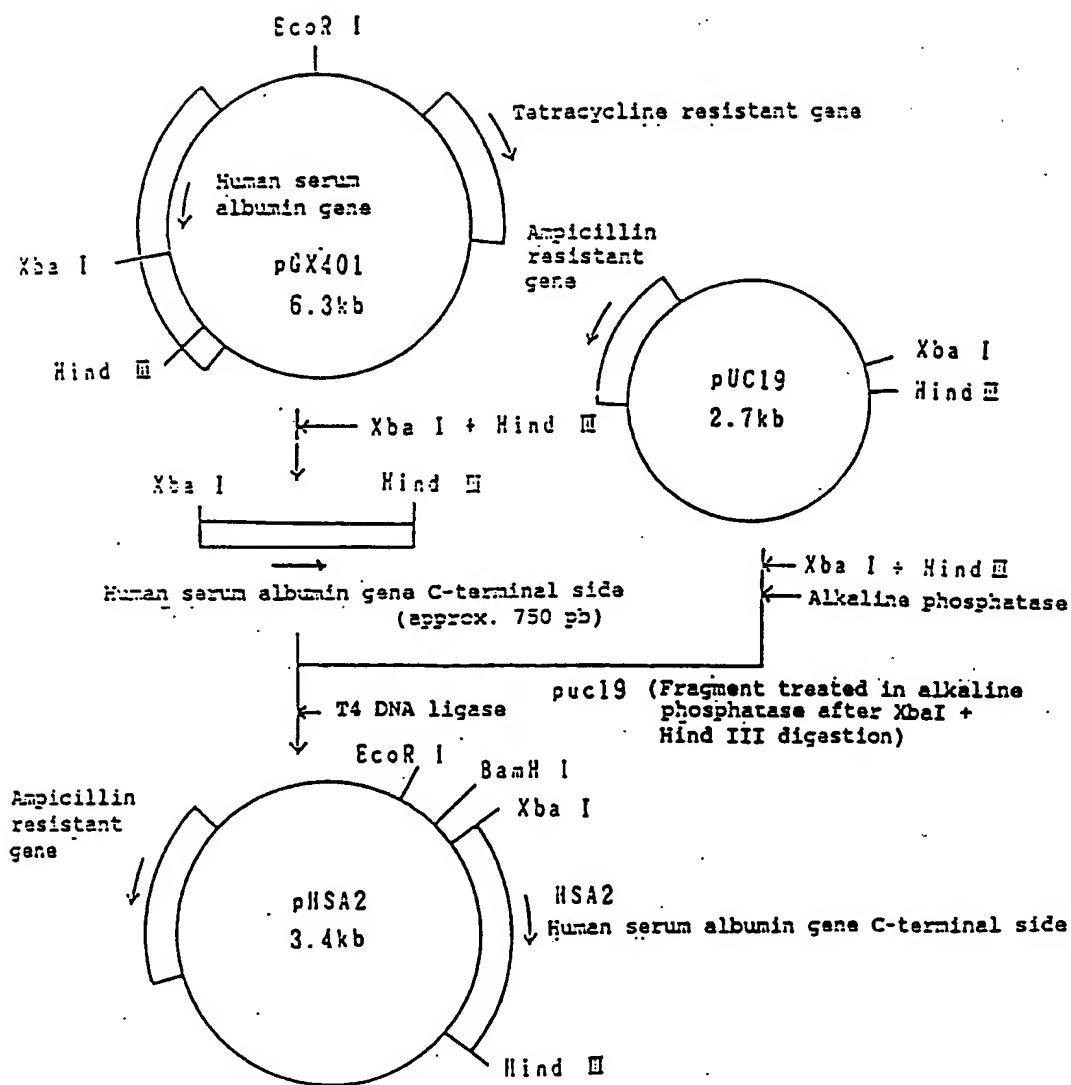
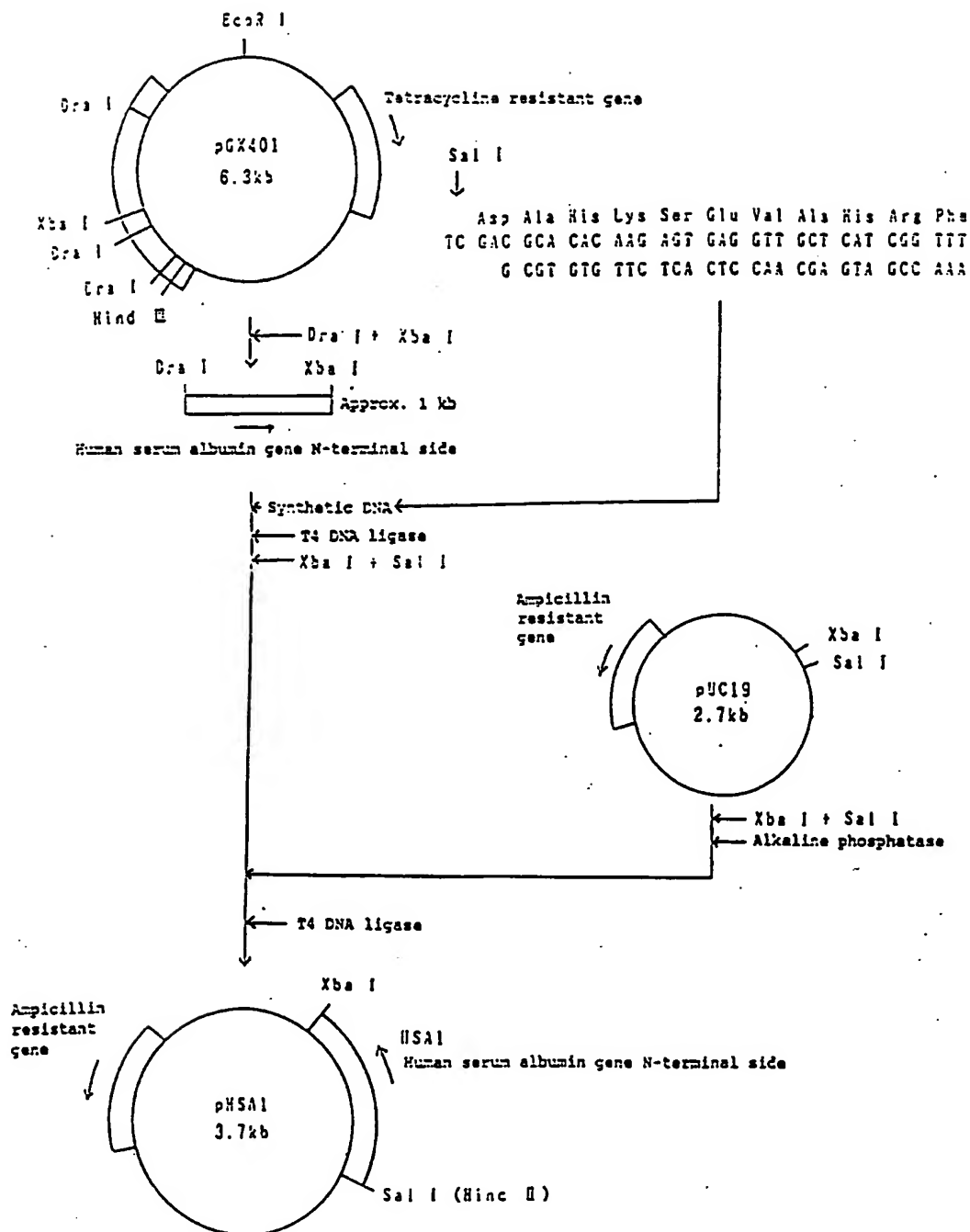


Fig. 7



Pl. 8

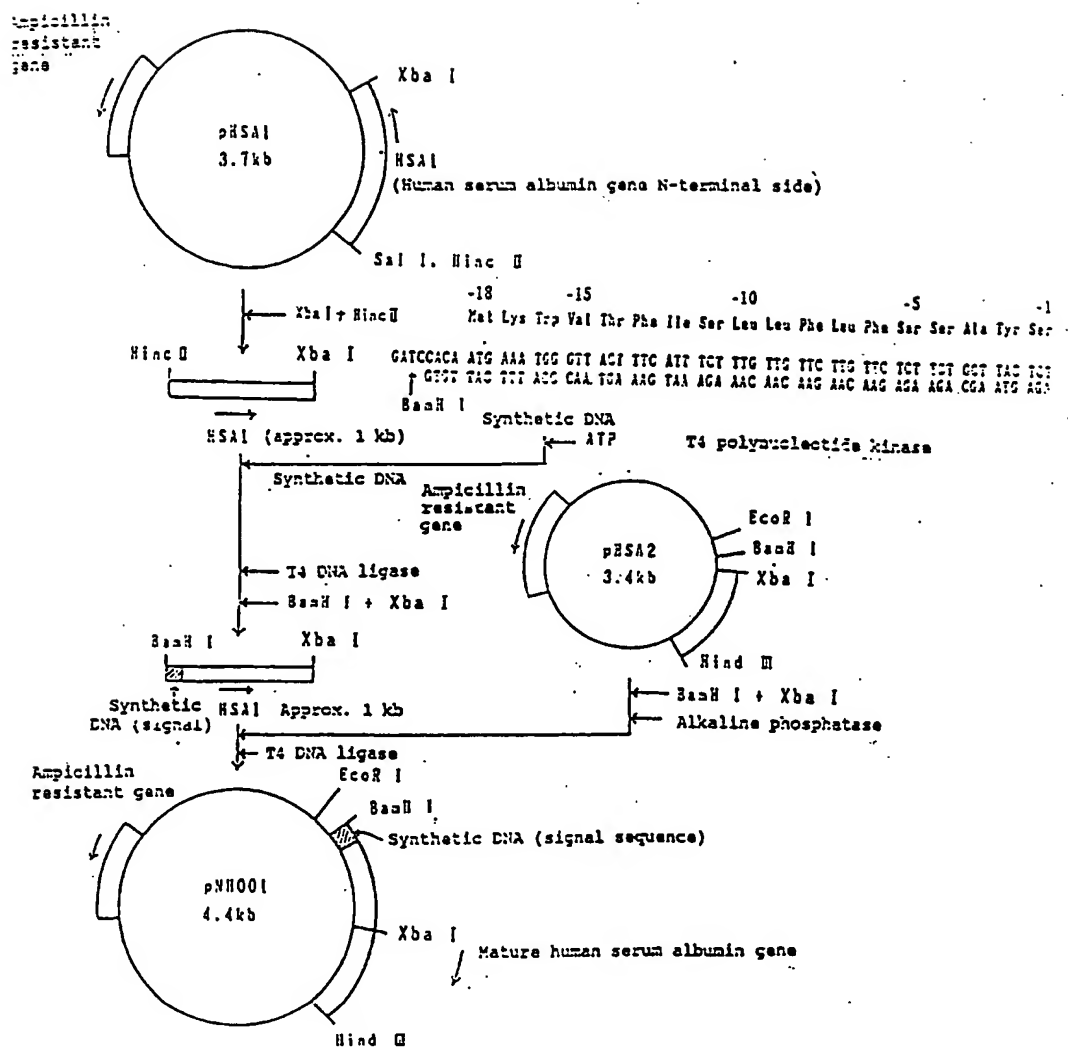


Fig. 9

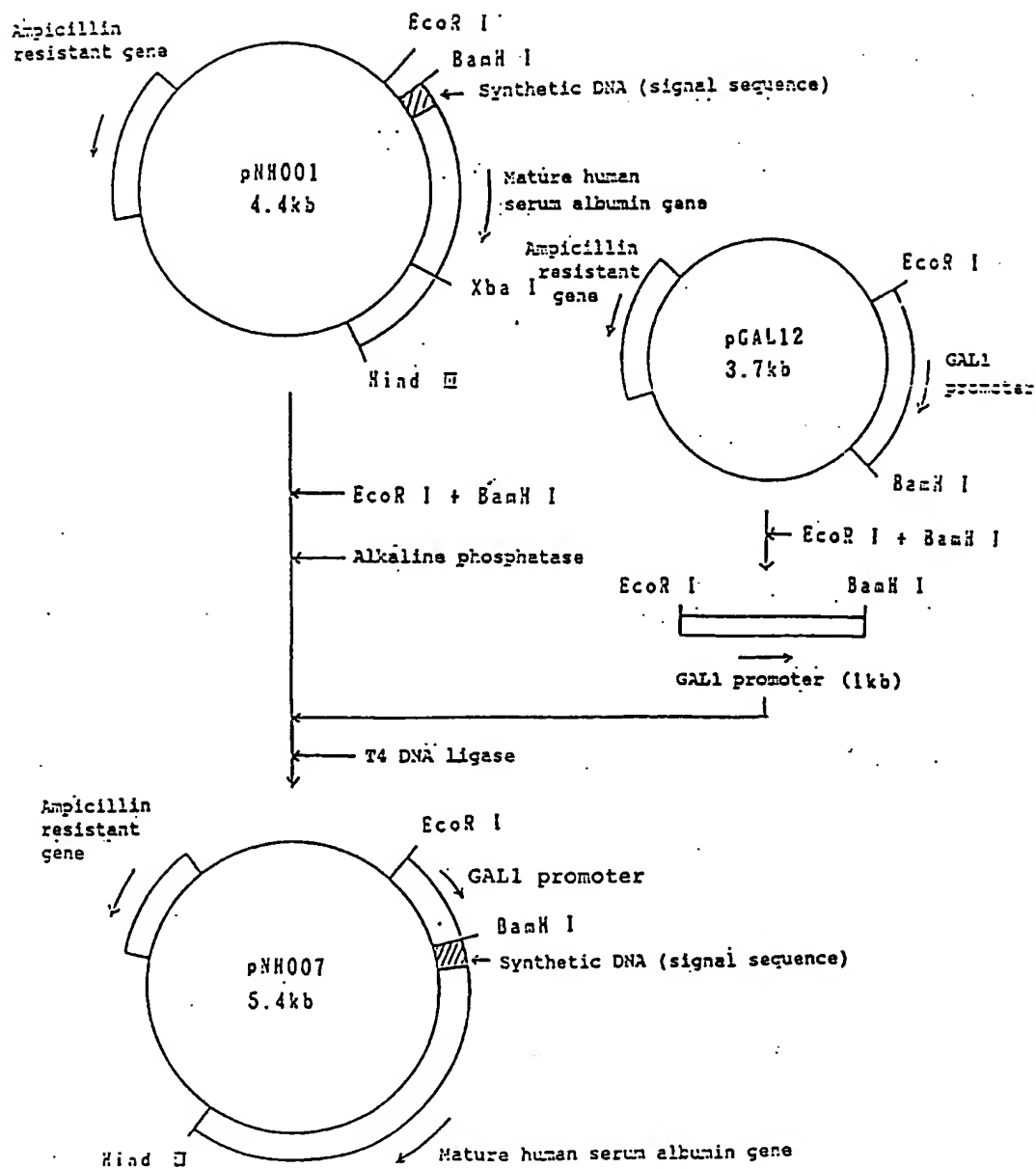
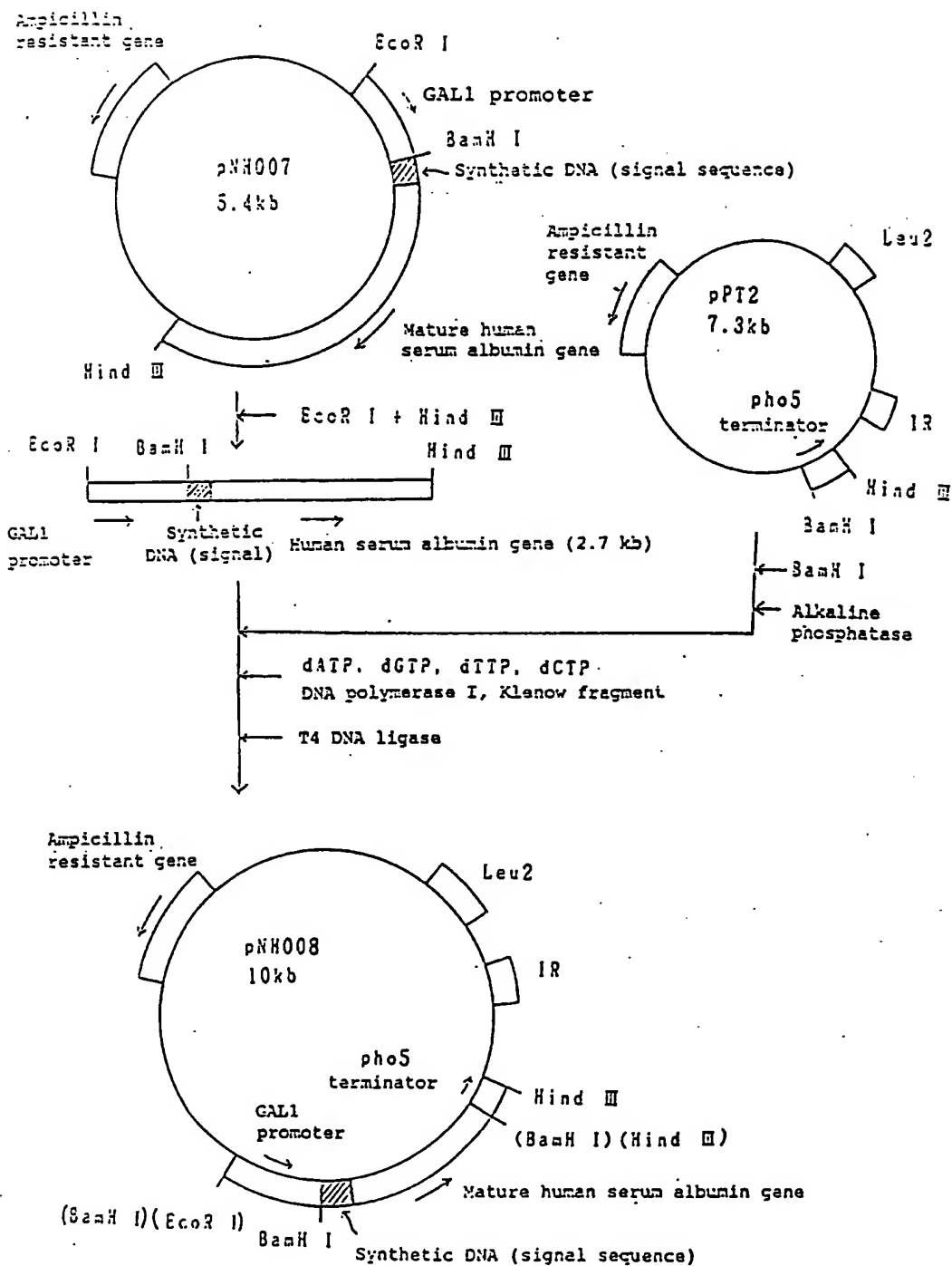


Fig. 10



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